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APPLICATION NO.	FILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCKET NO.	CONFIRMATION NO.
09/856,803	05/25/2001	Stephen B. Liggett	MWH-0029US	3706
25106 75	590 09/20/2002			
GENAISSANCE PHARMACEUTICALS 5 SCIENCE PARK NEW HAVEN, CT 06511			EXAMINER	
			MYERS, CARLA J	
			ART UNIT	PAPER NUMBER
			1634	11
			DATE MAILED: 09/20/2002	11

Please find below and/or attached an Office communication concerning this application or proceeding.

	- 1	Application No.	Applicant(s)		
Office Action Summary		09/856,803	LIGGETT, STEPHEN B.		
		Examiner	Art Unit		
		Carla Myers	1634		
	The MAILING DATE of this communic		th the correspondence address		
Period fo		D DEDLY IS SET TO EXPIDE 4 M	ONTUKO EBOM		
THE I - Exter after - If the - If NO - Failur - Any r	DRTENED STATUTORY PERIOD FO. MALING DATE OF THIS COMMUNIC issues of time ray, be available under the provisions of SN (6) MONTS's SN, or all shall be under the provisions of SN (6) MONTS's Specified above is less than they (30) period for raply specified above is less than they (30) period for raply a specified above in the maximum statu- te to raply within the set or extended period for raply will period for raply to office later than three months after dipatent term adjustment. See 37 CFR 1,704(b).	ATION.  37 CFR 1.136(a). In no event, however, may a relation.  days, a reply within the statutory minimum of thirt tory period will apply and will expire SIX (6) MON.  It by statute, cause the application to become AB.	eply be timely filed  y (30) days will be considered timely.  THS from the mailing date of this communication.  ANDONED (35 U.S.C. S 133).		
1)	Responsive to communication(s) filed	d on			
2a)		b)☐ This action is non-final.			
3)	Since this application is in condition f	,	ters, prosecution as to the merits is		
-	closed in accordance with the practic on of Claims				
4)⊠	Claim(s) <u>1-21,23 and 26-29</u> is/are per	nding in the application.			
	4a) Of the above claim(s) is/are	withdrawn from consideration.			
5)	Claim(s) is/are allowed.				
6)	Claim(s) is/are rejected.				
7)	Claim(s) is/are objected to.				
	Claim(s) <u>1-21,23 and 26-29</u> are subject	ct to restriction and/or election requ	irement.		
	on Papers				
9) 🔲 -	The specification is objected to by the I	Examiner.			
10)	The drawing(s) filed on is/are: a	, ,			
	Applicant may not request that any object				
11) ☐ The proposed drawing correction filed on is: a) ☐ approved b) ☐ disapproved by the Examiner.					
If approved, corrected drawings are required in reply to this Office action.					
	The oath or declaration is objected to b	y the Examiner.			
Priority u	nder 35 U.S.C. §§ 119 and 120				
13)	Acknowledgment is made of a claim for	or foreign priority under 35 U.S.C. §	§ 119(a)-(d) or (f).		
a)[	☐ All b)☐ Some * c)☐ None of:				
	<ol> <li>Certified copies of the priority do</li> </ol>	ocuments have been received.			
2. Certified copies of the priority documents have been received in Application No					
* S	Copies of the certified copies of application from the Internat ee the attached detailed Office action	tional Bureau (PCT Rule 17.2(a)).			
	cknowledgment is made of a claim for	•			
a	The translation of the foreign languages	uage provisional application has be	een received.		
Attachment		phone, and 50 0.0.0.	33		
1) Notice	e of References Cited (PTO-892) e of Draftsperson's Patent Drawing Review (PTC nation Disclosure Statement(s) (PTO-1449) Pap	0-948) 5) Notice of I	Summary (PTO-413) Paper No(s) nformal Patent Application (PTO-152)		

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- 1. Restriction to one of the following inventions is required under 35 U.S.C. § 121 and 372:
- I. Claims 1-8 and 11, drawn to a method of genotyping the  $\beta_2AR$  gene by determining the identity of a nucleotide in the 5' leader cistron of the  $\beta_2AR$  gene.
  - II. Claims 9-10 and 12, drawn to a method for detecting a 5' β<sub>2</sub>AR 5' LC peptide variant.
  - III. Claims 13-15, drawn to allele specific probes.
- $IV. \ Claims \ 16\text{-}17, \ drawn to methods for haplotyping by detecting the presence of 2 or more polymorphisms.$
- V. Claims 18, 28 and 29, drawn to methods for predicting a genotype in a coding sequence by detecting a polymorphism in the 5' leader cistron of the  $\beta_3$ AR gene.
- VI. Claims 19 and 20, drawn to a method for determining the frequency of a genotype of the  $\beta_2AR$  gene in a population.
- VII. Claims 19 and 20, drawn to a method for determining the frequency of a haplotype of the  $\beta_2AR$  gene in a population.
- VIII. Claims 21 and 23, drawn to a method for determining an association between a  $\beta_2AR$  gene polymorphism and a trait.
  - IX. Claims 26 and 27, drawn to a method of predicting a patients response to an agonist.
- 2. The inventions listed above do not relate to a single general inventive concept under PCT Rule 13.1 because, under PCT Rule 13.2, they lack the same or corresponding special technical feature for the following reasons:

The first claimed invention, claims 1-8 and 11, lacks unity because the invention represents methods which do not provide a special technical feature over the art. Claims 1-8 and 11 are broadly drawn to methods for genotyping the  $\beta_2AR$  gene by sequencing the 5' cistron leader of the  $\beta_2AR$  gene. Since the 5' cistron leader of the  $\beta_2AR$  gene was known in the art at the time the invention was made, as was methods of sequencing the 5' cistron leader sequence (See: Edmorine et al, Proceedings of the National Academy of Science, USA (1987) 84: 6995-6999),

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there is no special technical feature linking the recited groups, as would be necessary to fulfill the requirement for unity of invention.

Each of the claimed methods is patentably distinct from the other because they involve different method steps and have different objectives. The methods of invention I require steps of determining the sequence in the 5' cistron leader of the  $\beta_2AR$  gene to achieve the objective of genotyping the  $\beta_2AR$  gene. The methods of invention II require analyzing the amino acid sequence of the  $\beta_2AR$  gene 5' cistron leader in order to accomplish the objective of detecting a  $\beta_2AR$  variant peptide. The methods of invention IV require identifying the nucleotide sequence at 2 or more positions in the  $\beta_2AR$  gene in order to achieve the objective of haplotying the  $\beta_2AR$ gene. The methods of invention V require predicting a genotype in the coding sequence of the β<sub>2</sub>AR gene by detecting a genotype of the 5' leader cistron. Invention VI involves determining the frequency of a genotype of the  $\beta_2$ AR gene in the population. Invention VII involves analyzing a population to determine the frequency of a β<sub>2</sub>AR gene haplotype. Invention VIII involves comparing the frequency of a  $\beta_2AR$  gene polymorphism in a reference population and in a population exhibiting a trait in order to identify an association between the  $\beta_2AR$  gene polymorphism and the trait. Invention IX involves detecting a genotype in the β<sub>2</sub>AR gene 5' leader cistron in order to predict a patients response to an agonist of  $\beta_2AR$ . Further, each of the claimed methods are distinct over the allele specific probes because the allele specific probes can be used in patentably distinct methods, such as methods for detecting the β<sub>2</sub>AR gene. 3.

## Sequence and Polymorphism Election

In addition, Inventions I and III detailed above read on patentably distinct inventions drawn to nucleic acids comprising patentably distinct sequences. The nucleotide sequences are patentably distinct because they are structurally and functionally unrelated sequences, and a further restriction is applied to each invention. A search of one nucleotide sequence would not lead one of skill in the art to each of the other nucleotide sequences and thereby distinct searches

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are required for each of the nucleotide sequence and the searches are not coextensive. In response to the restriction requirement, Applicant must further elect a single polynucleotide selected from the group consisting of SEQ ID NO: 5-10.

It is noted that nucleotide sequences encoding different proteins and/or having distinct nucleotide sequences are structurally distinct chemical compounds and are unrelated to one another. These sequences are thus deemed to constitute independent and distinct inventions. Absent evidence to the contrary, each such nucleotide sequence is presumed to represent an independent and distinct invention, subject to a restriction requirement.

Furthermore, Applicants must elect a single polymorphism to be examined. Each alteration in the sequence of the  $\beta_2AR$  gene is unique and unobvious over the other. The haplotypes and genotypes encompassed by these claims are distinct from each other and from the single polymorphisms. For example, a molecule of haplotype 1, comprising a particular combination of polymorphisms, differs chemically, structurally, and functionally from a molecule of haplotype 2 and from a molecule comprising a single polymorphism. The special technical feature of each haplotype or genotype is the combination of polymorphisms contained therein, which feature is lacking from and not shared with each other haplotype or genotype or with, e.g., a molecule comprising any single polymorphism set forth in the claims. Further, a search for references teaching a polymorphism at position -20PS would not lead one of skill in the art to references teaching a polymorphism at position +46PS. For an elected invention that involves genotyping, Applicant is required to elect one of the polymorphisms selected from the group consisting of -20PS, +46PS, +79PS, +100PS and +491PS. With respect to inventions I and III, the elected polymorphism must correspond to the elected allele specific polynucleotide sequence (SEQ ID NO: 5-10). For an elected invention that involves haplotyping, Applicant is required to elect a single haplotype to be examined.



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Applicant is advised that the reply to this requirement to be complete must include an election of the invention to be examined even though the requirement be traversed (37 CFR 1.143).

Any inquiry concerning this communication or earlier communications from the examiner should be directed to Carla Myers whose telephone number is (703) 308-2199. The examiner can normally be reached on Monday-Thursday from 6:30 AM-5:00 PM.

If attempts to reach the examiner by telephone are unsuccessful, the examiner's supervisor, W. Gary Jones, can be reached on (703)-308-1152. The fax number for the Technology Center is (703)-305-3014 or (703)-305-4242.

Any inquiry of a general nature or relating to the status of this application should be directed to the receptionist whose telephone number is (703) 308-0196.

Carla Myers

September 12, 2002

CARLA J. MYERS